

## SEQUENCE LISTING

<110>	Ingvar M. Ferby et al.
<120>	PROTEIN WITH CELL PROLIFERATION AND CELL DIVISION MODULATING ACTIVITY AND DNA ENCODING SUCH PROTEIN
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<140> <141>	09/889,592 .:001-08-02
	PCT/EP00/00877 2000-02-03
	EP 99102172.6 1999-02-03
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atcaac	octs gggseggtgt eddectttet asa atg agg cac atg sag agt gta 234 Met Arg His Met Gln Ser Val 1 5
_	g geo age toc att tgt ggo age ygg gtg aag bag gtb att ggo ——282 g Ala Ser Ser Ile Cys Gly Ser Gly Val Lys Gln Val Ile Gly ————————————————————————————————————
	g dat dog dad god ogg gtt gtt gga gog ogd åan gly fallal i i i i i i i i i i i i i i i i i
	g aga gag gag tig toa gio aaa ooc aaa aig gig oga aai acc 378 u Arg Glu Gli Leu Ser Val Lys Pro Lys Mei Val Arg Asn Thr 45 50 55
cat ct	c aat ota dag doo dag gag ogd dag god tto tad agg otd ota 426

His Leu Asn Leu Gln Pro Gln Glu Arg Gln Ala Phe Tyr Arg Leu Leu 60 gaa aat gag cag att cag gaa ttc ctt tct atg gac tcc tgt cta agg 474 Glu Asn Glu Gln Ile Gln Glu Phe Leu Ser Met Asp Ser Cys Leu Arg att too gao aag tat oto ata goa atg gtt ota goa tat tit aag ogg 522 Ile Ser Asp Lys Tyr Leu Ile Ala Met Val Leu Ala Tyr Phe Lys Arg 95 90 570 gea geg gge etc tac acc age gag tac aca acc atg aat tte ttt gtt Ala Ala Gly Leu Tyr Thr Ser Glu Tyr Thr Thr Met Asn Phe Phe Val 105 gee etg tat etg get aat gae atg gag gaa gat gaa gae tat aaa 618 Ala Leu Tyr Leu Ala Asn Asp Met Glu Glu Asp Glu Glu Asp Tyr Lys 125 tat gaa ato tto coo tgg goa cta gga gao tog tgg cgt gag ott tto 666 Tyr Glu Ile Phe Pro Trp Ala Leu Gly Asp Ser Trp Arg Glu Leu Phe 140 cca caa ttt ttg cgt ctc cgg gac gac ttc tgg gct aaa atg aac tac 714 Pro Gln Phe Leu Arg Leu Arg Asp Asp Phe Trp Ala Lys Met Asn Tyr 160 ega gdwigtt. 31' agr ogn add tod tot gal dag gta atd tod aaa gat 762 Arg Ala Val Val Ser Arg Arg Cys Cys Asp Glu Vai Met Ser Lys Asp 175 ccc act cat tgg gcc tgg ctg aga gat cgc ccc atg cat cac agc ggg 810 Pro Thr His Trp Ala Trp Leu Arg Asp Arg Pro Met His His Ser Gly 190 858 ged atg cgt ggt tad oft aga aad gag gad gad tit tid odd ogg ggt Ala Met Arg Gly Tyr Leu Arg Asn Glu Asp Asp Phe Phe Pro Arg Gly 205 200 cca ggd ctt aca cca ged agd tgt aca ctt tgd cat aaa gca ggt gtd 906 Pro Gly Leu Thr Pro Ala Ser Cys Thr Leu Cys His Lys Ala Gly Val tgt gad tot ggt ggg gto tod dad aad tot tod tot oca gaa daa 954 Cys Asp Ser Gly Gly Val Ser His Asn Asn Ser Ser Pro Glu Gln 240 235 yay uti tit han han ann aat agg gag tee eag gag ett ete atg 1002 Glu Ile Phe His Tyr Thr Asn Arg Glu Trp Ser Jin Giu Del Le. Met 250 ting our cot gay etg ting ont gat and gag tigt act cat gad tha cad 1050 Leu Pro Pro Glu Leu Leu Asp Pro Glu Cys Thr His Asp Leu His 270 265 att etc cag gag eca ttg gtt gga tta gag eca gat ggg acg geg etg 1098 Ile Leu Gln Glu Pro Leu Val Gly Leu Glu Pro Asp Gly Thr Ala Leu

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Gln Ala Phe Tyr	Arg Leu Leu Glu 70	Asn Glu Gin Ile Gi: 75	n Glu Phe Len 80
Ser Met Asp Ser	Cys Leu Arg 11e 85	ser Asp Lys Tyr Lon 90	tile Ala Met 95
Val Leu Ala Tyr 100	Phe Lys Arg Ala	Ala Gly Leu Tyr Th	r Ser Glu Tyr 110

Thr Thr Met Asn Phe Phe Val Ala Leu Tyr Leu Ala Asn Asp Met Giu 120 Glu Asp Glu Glu Asp Tyr Lys Tyr Glu Lie Phe Pro Tip Ala Leu Gly 135 140 Asp Ser Trp Arg Glu Leu Phe Pro Gln Phe Leu Arg Leu Arg Asp Asp 145 150 155 160 Phe Trp Ala Lys Met Asn Tyr Arg Ala Val Val Ser Arg Arg Cys Cys Asp Glu Val Met Ser Lys Asp Pro Thr His Trp Ala Trp Leu Arg Asp Arg Pro Met His His Ser Gly Ala Met Arg Gly Tyr Leu Arg Asn Glu Asp Asp Phe Phe Pro Arg Gly Pro Gly Leu Thr Pro Ala Ser Cys Thr 215 Leu Cys His Lys Ala Gly Val Cys Asp Ser Gly Gly Val Ser His Asn 230 235 Asn Ser Ser Pro Glu Gln Glu Ile Phe His Tyr Thr Asn Arg Glu 245 250 255 Trp Ser Gln Glu Leu Leu Met Leu Pro Pro Glu Leu Leu Leu Asp Pro 260 265 270 Glu Cys Thr His Asp Leu His Ile Leu Gln Glu Pro Leu Val Gly Leu 275 280 285 Glu Pro Asp Gly Thr Ala Leu Glu Trp His His Leu 290 295 300 <210 > 3 -:211: 1357

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tttgotoaga toaaoootog gttogttgto occotttota oa atg agg oat atg Met Arg His Met 1	174
cag agt gea acc egg gec acc tta gtt tgt ggc age ggg gta aag cag Gin Ser Ala Thr Arg Ala Thr Leu Val Cys Gly Ser Gly Val Lys Gln 5 10 15 20	22.1
ate att ged aag gga dat eeg aat aed egg gtt tit gga geg egd aag Ile Ile Ala Lys Gly His Pro Asn Thr Arg Val Phe Gly Ala Arg Lys 25 30 35	27.)
gog aaa ato oot gag aga gag gtg ota god god aaa ooc aag ato aog Ala Lys Ile Pro Glu Arg Glu Val Leu Ala Ala Lys Pro Lys Ile Thr 40 45 50	318
ogo att aca cat etc dat eta caa dec bag gag ogo dag god tit tad Ang Ile Thr His Leu Ash Leu Gln Pro Gln Glu Ang Gin Ala Phe Tyr 67 63	364
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ttt fto one hae fff fta ogt oto ogg gad gad tto ogg got aaa åtg Phe Phe Pro Gin Phe Leu Arg Leu Arg Asp Asp Pho Trp Ala Bys Met 130 - 155 - 160	654
Asn Tyr Arg Ala Val Val Ser Arg Arg Cys Cys Asp Glu Val Met Ala 165 170 175 180	702

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Leu Glu Trp His His Leu 295	
•	1146
29 <sup>5</sup>	1146 1206
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260 265 270

Thr Tyr Asp Ile His Ile Phe Gln Glu Pro Leu Val Gly Leu Glu Pro 275 280 285

Asp Gly Ala Ala Leu Glu Trp His His Leu 290 295